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Applicant(s): Michael R. Green et al.
ALTERING VIRAL TROPISM
REPLACEMENT SHEET

Fig. 8

Moloney Murine Leukemia Virus – envelope protein (gp70), nucleic acid sequence (from complete MoMLV genome sequence; Genbank Accession No. NC\_001501). The SU (extracellular domain) is encoded by nucleotides 5612 - 6919 (pictured below). The transmembrane and cytoplasmic tail extends from nucleotides 6920-7507. There is a signal peptide sequence at the beginning of the SU, localizing the protein to the cell membrane.

aattcttctg	atgctcagag	gggtcagtac	tgcttcgccc	ggctccagtc	ctcatcaagt	5581
ctataatatc	acctgggagg	taaccaatgg	agatcgggag	acggtatggg	caacttctgg	5641
caaccaccct	ctgtggacct	ggtggcctga	ccttacccca	gatttatgta	tgttagccca	5701
ccatggacca	tcttattggg	ggctagaata	tcaatcccct	ttttcttctc	ccccggggcc	5761
cccttgttgc	tcagggggca	gcagcccagg	ctgttccaga	gactgcgaag	aacctttaac	5821
ctccctcacc	cctcggtgca	acactgcctg	gaacagactc	aagctagacc	agacaactca	5881
taaatcaaat	gagggatttt	atgtttgccc	cgggccccac	cgcccccgag	aatccaagtc	5941
atgtgggggt	ccagactcct	tctactgtgc	ctattggggc	tgtgagacaa	ccggtagagc	6001
ttactggaag	ccctcctcat	catgggattt	catcacagta	aacaacaatc	tcacctctga	6061
ccaggctgtc	caggtatgca	aagataataa	gtggtgcaac	cccttagtta	ttcggtttac	6121
agacgccggg	agacgggtta	cttcctggac	cacaggacat	tactggggct	tacgtttgta	6181
tgtctccgga	caagatccag	ggcttacatt	tgggatccga	ctcagatacc	aaaatctagg	6241
		caaaccccgt				6301
		cagtcaccaa				6361
		cggaaaatag				6421
		ctgacaaaac				6481
		ttgccgtcct				6541
		cccaacacaa				6601
		ccaaaacaca				6661
		tagttgcccc				6721
gcttactcca	tgcatctcca	ccaccatact	gaaccttacc	actgattatt	gtgttcttgt	6781
cgaactctgg	ccaagagtca	cctatcattc	ccccagctat	gtttacggcc	tgtttgagag	6841
atccaaccga	cacaaaagag	aaccggtgtc	gttaaccctg	gccctattat	tgggtggact	6901
aaccatgggg	ggaattgccg	ctggaatagg	aacagggact	actgctctaa	tggccactca	6961
gcaattccag	cagctccaag	ccgcagtaca	ggatgatctc	agggaggttg	aaaaatcaat	7021
ctctaaccta	gaaaagtctc	tcacttccct	gtctgaagtt	gtcctacaga	atcgaagggg	7081
cctagacttg	ttatttctaa	aagaaggagg	gctgtgtgct	gctctaaaag	aagaatgttg	7141
cttctatgcg	gaccacacag	gactagtgag	agacagcatg	gccaaattga	gagagaggct	7201
taatcagaga	cagaaactgt	ttgagtcaac	tcaaggatgg	tttgagggac	tgtttaacag	7261
atccccttgg	tttaccacct	tgatatctac	cattatggga	cccctcattg	tactcctaat	7321
gattttgctc	ttcggaccct	gcattcttaa	tcgattagtc	caatttgtta	aagacaggat	7381
		ttttgactca				7441
cgagccatag	ataaaataaa	agattttatt	tagtctccag	aaaaaggggg	gaatgaaaga	7501